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FIG. 6

PROCESS	WORLD WIDE WEB SITE PAGE
sequence format conversion	<a href="http://dot.imgen.bcm.tmc.edu:9331/seq-util/Options/-feedseq.html">http://dot.imgen.bcm.tmc.edu:9331/seq-util/Options/-feedseq.html</a>
reverse complementation	<a href="http://dot.imgen.bcm.tmc.edu:9331/seq-util/Options/revecomp.html">http://dot.imgen.bcm.tmc.edu:9331/seq-util/Options/revecomp.html</a>
sequence translation	<a href="http://dot.imgen.bcm.tmc.edu:9331/seq-util/Options/sixframe.html">http://dot.imgen.bcm.tmc.edu:9331/seq-util/Options/sixframe.html</a>
protein information	<a href="http://www.expasy.ch/tools">http://www.expasy.ch/tools</a> ExPASy Proteomics Tools
sequence alignments using Clustal W	<a href="http://dot.imgen.bcm.tmc.edu:9331/multi-align/Options/elustaln.html">http://dot.imgen.bcm.tmc.edu:9331/multi-align/Options/elustaln.html</a>
sequence comparisons using BLAST 2.0	<a href="http://www.ncbi.nlm.nih.gov/gorf/bf2.html">http://www.ncbi.nlm.nih.gov/gorf/bf2.html</a> National Center For Biotechnology
sequence searches using BLAST 2.0	<a href="http://www.ncbi.nlm.nih.gov/blast/blast.cgi?Jform=0">http://www.ncbi.nlm.nih.gov/blast/blast.cgi?Jform=0</a> National Center for Biotechnology